

Reconciling qualitative and abstract (and scalable) modelling of biological regulatory networks

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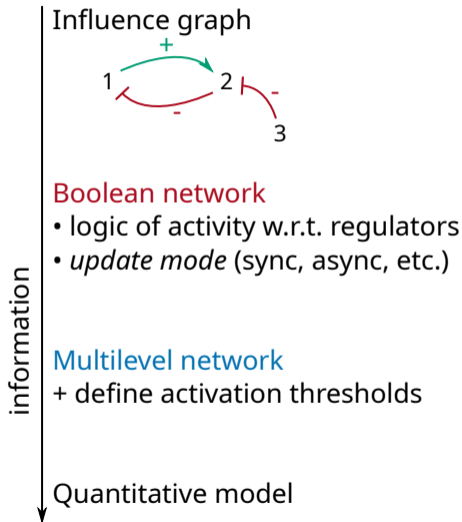
<https://loicpauleve.name>

joint work with

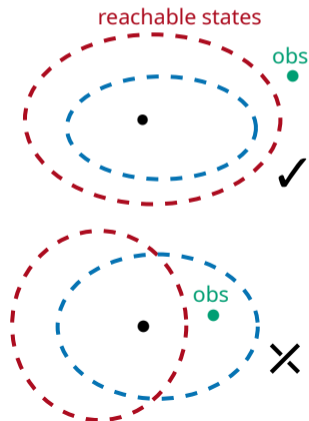
- T. Chatain, S. Haar, J. Kolčák (LSV/Inria, ENS Cachan, France)



Qualitative vs abstract modelling



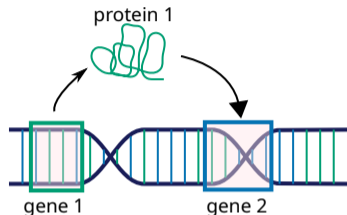
Consistency
analysis at Boolean level
transposable to multilevel?



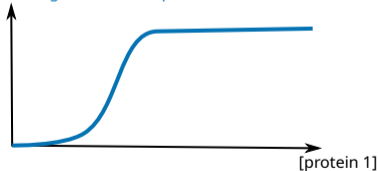
Most permissive semantics of Boolean networks

Gene expression is not Boolean

Qualitative modelling: Boolean vs multivalued networks

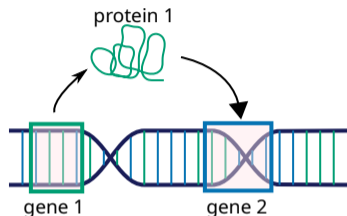


effect on gene 2 transcription



Gene expression is not Boolean

Qualitative modelling: Boolean vs multivalued networks

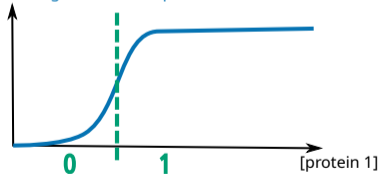


Boolean network

$$f : \mathbb{B}^n \rightarrow \mathbb{B}^n$$

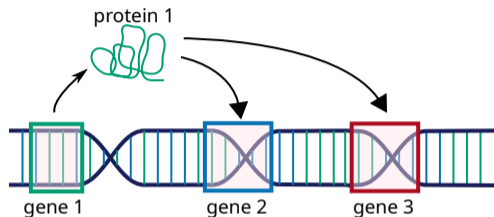
$$f_2(x) \triangleq x_1$$

effect on gene 2 transcription



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Qualitative modelling: Boolean vs multivalued networks



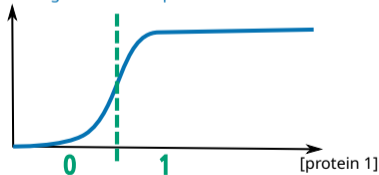
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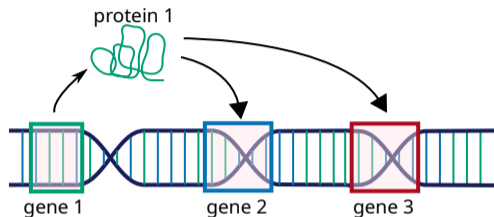
$$f_3(x) \triangleq x_1$$

effect on gene 2 transcription

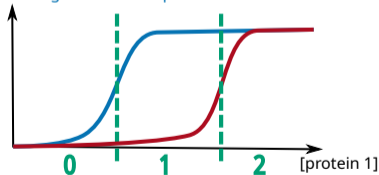


Gene expression is not Boolean

Qualitative modelling: Boolean vs multivalued networks



effect on gene 3 transcription
effect on gene 2 transcription



Boolean network

$$f : \mathbb{B}^n \rightarrow \mathbb{B}^n$$

$$f_2(x) \triangleq x_1$$

$$f_3(x) \triangleq x_1$$

Multivalued network

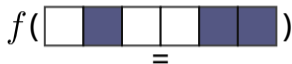
$$F_2(x) \triangleq (x_1 \geq 1)$$

$$F_3(x) \triangleq (x_1 \geq 2)$$

Introduction

Boolean Network (BN) $f : \mathbb{B}^n \rightarrow \mathbb{B}^n$

Configuration: $x \in \mathbb{B}^n$



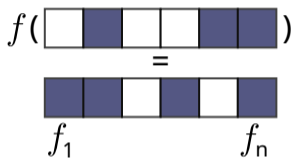
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↓ **synchronous** update



Introduction

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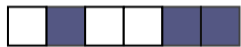
synchronous update



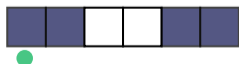
$$f\left(\begin{array}{|c|c|c|c|c|c|} \hline \square & \blacksquare & \square & \square & \blacksquare & \blacksquare \\ \hline \end{array}\right) =$$

$$\begin{array}{|c|c|c|c|c|c|} \hline \blacksquare & \blacksquare & \square & \blacksquare & \square & \blacksquare \\ \hline \end{array}$$

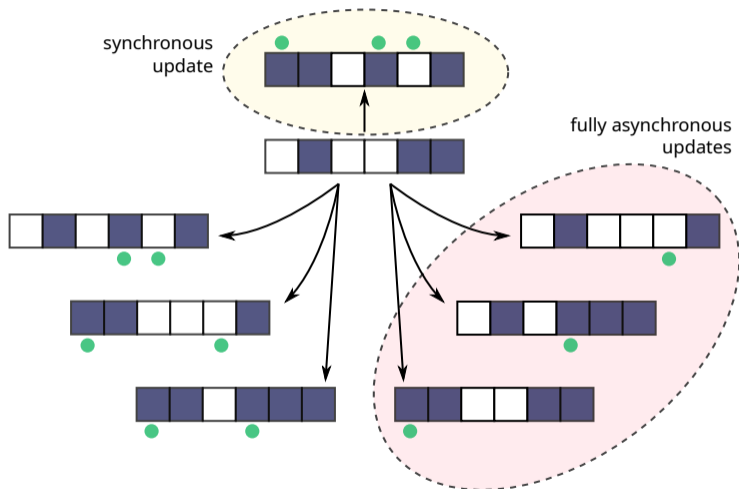
$f_1 \qquad \qquad \qquad f_n$



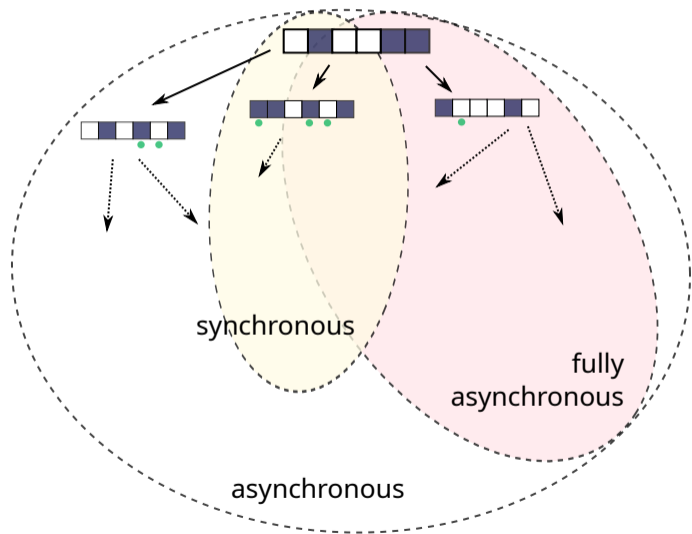
fully asynchronous updates



Asynchronous updates



Reachable configurations



Asynchronous
Boolean networks:
inconsistent abstraction

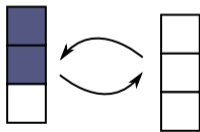
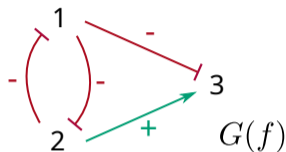
Motivating example

(embedded in many actual biological networks)

$$f_1(x) \triangleq \neg x_2$$

$$f_2(x) \triangleq \neg x_1$$

$$f_3(x) \triangleq \neg x_1 \wedge x_2$$



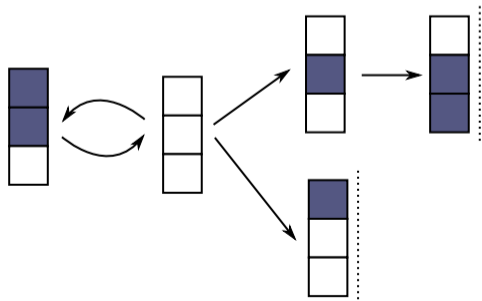
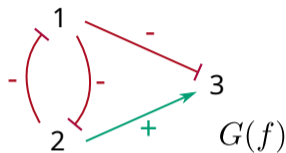
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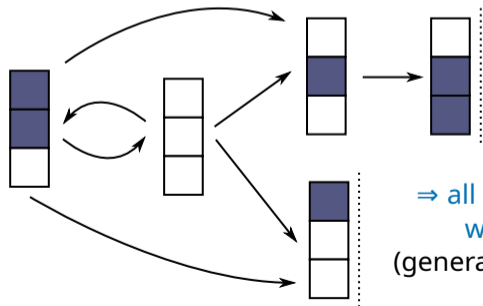
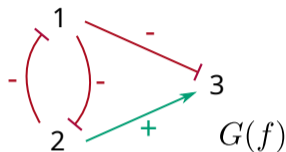
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⇒ all configurations reachable
with any update mode
(generalized) asynchronous mode

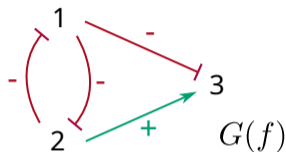
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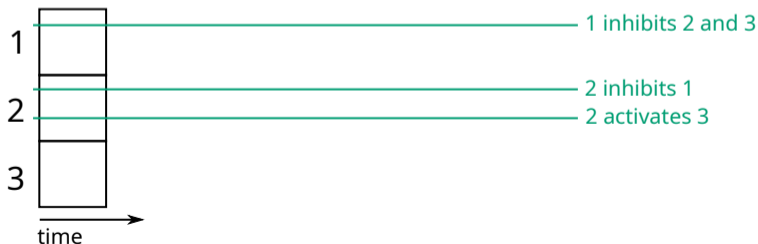
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Compatible **continuous/multilevel** model:



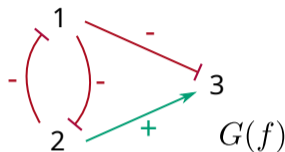
Motivating example

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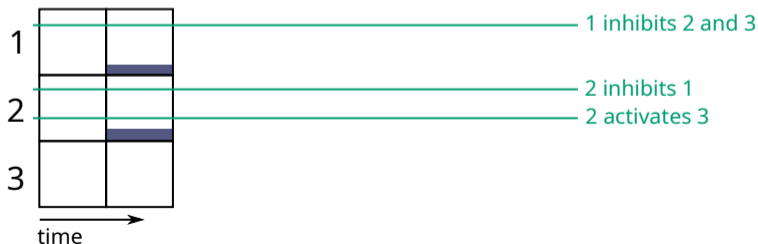
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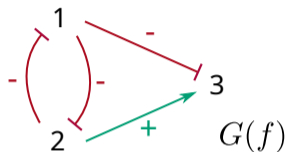
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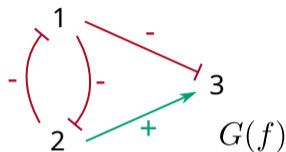
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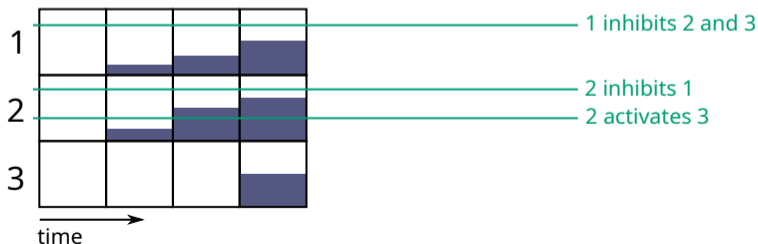
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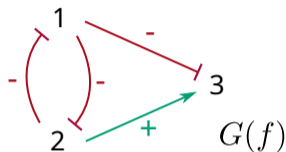
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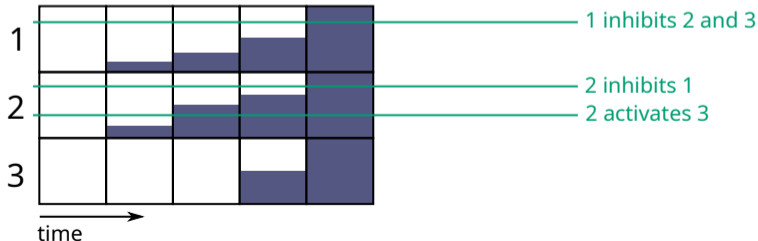
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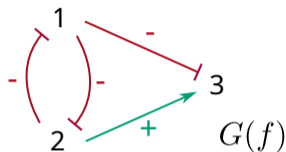
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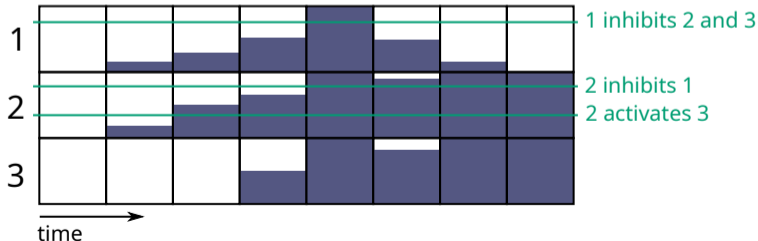
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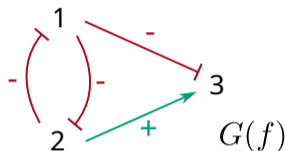
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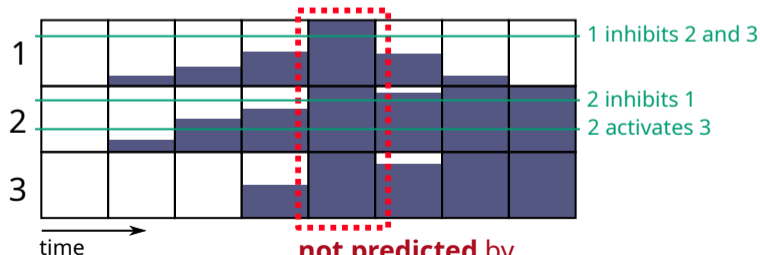
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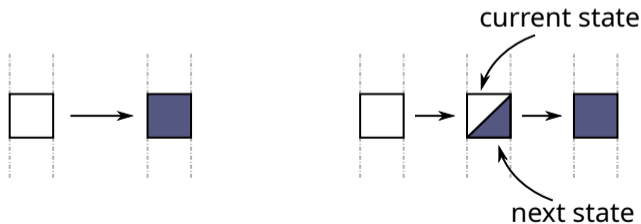


**not predicted by
update modes in Boolean**

Most Permissive
Boolean networks
enabling new behaviours

Most permissive semantics

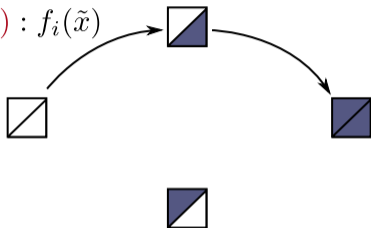
- **delay between firing and application** of state change
⇒ allow interleaving other state changes
- in "intermediate" states 
other components choose what they see



Most permissive semantics

Automaton of component i :

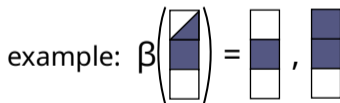
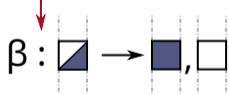
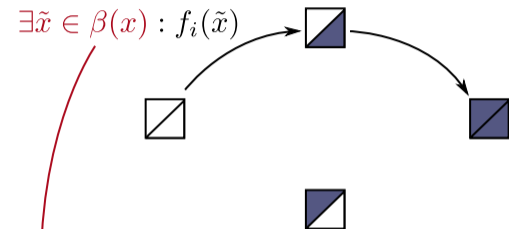
$$\exists \tilde{x} \in \beta(x) : f_i(\tilde{x})$$



+ fully-asynchronous interleaving

Most permissive semantics

Automaton of component i :

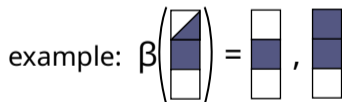
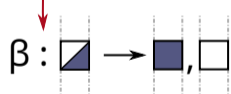
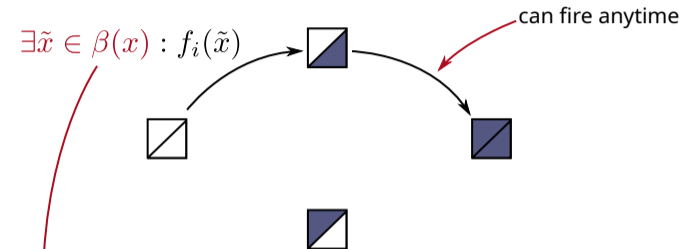


Choose value of "changing" components
(act as choosing an activation threshold)

+ fully-asynchronous interleaving

Most permissive semantics

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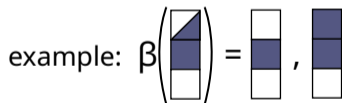
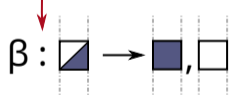
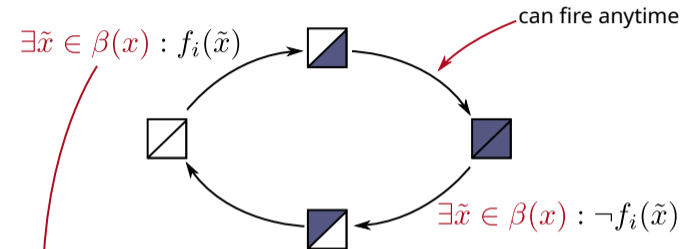


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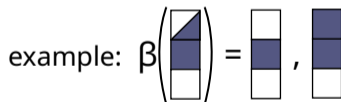
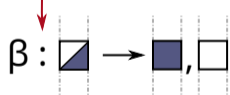
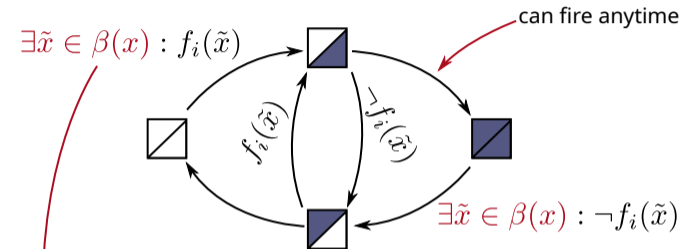


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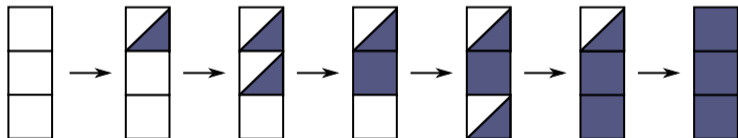
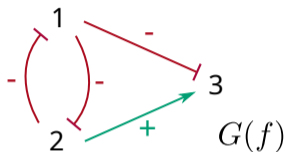
+ fully-asynchronous interleaving

Application to motivating example

$$f_1(x) \triangleq \neg x_2$$

$$f_2(x) \triangleq \neg x_1$$

$$f_3(x) \triangleq \neg x_1 \wedge x_2$$



Properties of the most permissive semantics

Correct abstraction of multilevel/quantitative systems:

- includes all the **transitions of every update mode**
- quantitative **refinements only remove behaviours**

- **Reachability** can be decided in **quadratic nb of transitions**
⇒ y is reachable from x iff there is a path of length at most $3n$
(PTIME with locally-monotonic networks, or encoded as BDDs/Petri nets/...;
NP-complete otherwise; instead of PSPACE-complete with update modes)

- **Attractors are hypercubes** (minimal trap spaces)
⇒ existence within a sub-space is **NP-complete**
(instead of PSPACE-complete)
⇒ fixpoints are the same as with update modes

Refinements of Boolean Networks

A **multivalued network**

$$F : \mathbb{M}^n \rightarrow \{-1, 0, 1\}^n$$

is a **refinement of a Boolean network** f iff

$$F_i(\begin{array}{|c|c|c|} \hline \square & \blacksquare & \square \\ \hline \end{array}) > 0 \implies \exists \begin{array}{|c|c|c|} \hline \text{---} & \blacksquare & \text{---} \\ \hline \end{array} : f_i(\begin{array}{|c|c|c|} \hline \square & \blacksquare & \square \\ \hline \end{array}) = 1$$

$$F_i(\begin{array}{|c|c|c|} \hline \blacksquare & \square & \square \\ \hline \end{array}) < 0 \implies \exists \begin{array}{|c|c|c|} \hline \blacksquare & \text{---} & \text{---} \\ \hline \end{array} : f_i(\begin{array}{|c|c|c|} \hline \blacksquare & \square & \square \\ \hline \end{array}) = 0$$

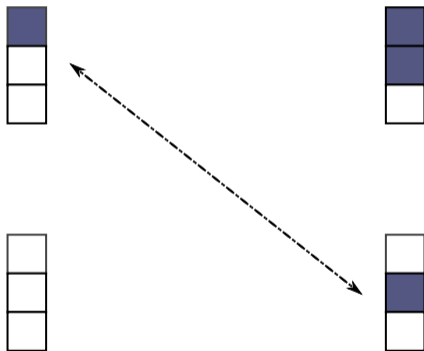
**Most permissive semantics weakly simulates
any multivalued refinement with *any update mode***

(can be extended to ODEs)

Attractors of the most permissive semantics

Attractor: smallest set of configurations closed by transitions

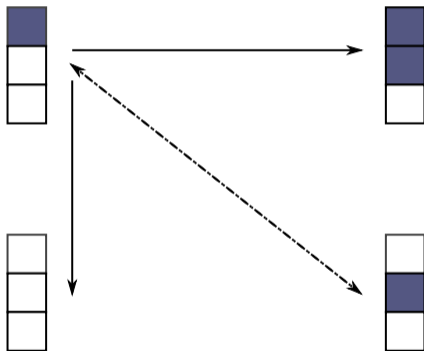
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Attractors of the most permissive semantics

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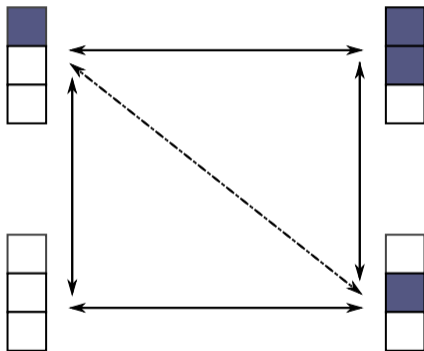
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Attractors of the most permissive semantics

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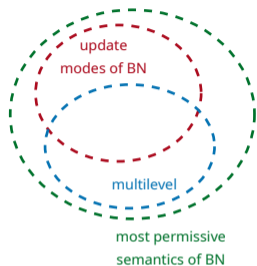
Attractors are hypercubes



Attractors of most permissive semantics = minimal trap spaces

Deciding if trap space is minimal = 2 reachability checking

Is most permissive semantics restrictive?



Minimality of abstraction

to any "most permissive" transition, there is corresponding multilevel transition
Work in progress w/ "most permissive" paths: non-minimal, but tricky counter-examples

- **fixpoints** are **preserved (identical)**
 - **trap spaces**: known to be relevant for reasoning with **attractors**
[Klarner et al in Nat. Comp. 2015] [Naldi in Front. Phys. 2018]
- ⇒ most permissive semantics seems still adequate to model **differentiation processes** !

Conclusion

Update modes of Boolean networks (sync, async, etc.):

- difficult to justify for biology (strong implications on dynamics)
 - can **miss important behaviours**
- ⇒ lead to **reject valid models** of biological systems...
- have limited tractability (model-checking, ...)

Most permissive semantics:

- **correct abstraction**: guarantees that multilevel/quantitative refinements only remove behaviours
 - **simpler complexity**: reachability PTIME, attractors NP
- ⇒ genome-scale tractability (BNs with >10,000 components)

Scalable Boolean analysis consistent with quantitative models